

*5 10 15 20 25 30 35 40 45 50 55 60 65 70
 GAATTOGGCA CGAGCAACA GTCATTAGTG GAGGACTCTA TTGTAATAAA CTGTGCTTTA AAATGTAAC
 75 80 85 90 95 100 105 110 115 120 125 130 135
 CAGGGAGOGT TTTTITTOCT CACATTGTCC AGAAGCAACC TTTCTTCTG AGOCTGGATT AATC ATG
 140 145 150 155 160 165 170 175 180 185 190
 AGA GAG CTC GTC AAC ATT CCA CTG TTA CAG ATG CTC ACC CTG GTT GGC TTC AGC GGG
 R E L V N I P L L Q H L T L V A F S G>
 195 200 205 210 215 220 225 230 235 240 245 250
 ACC GAG AAA CTT CCA AAA GGC OCT GTC ATC ACC ACG OCT CTT GAA ACT GTA GAT GGC
 T E K L P K A P V I T T P L E T V D A>
 255 260 265 270 275 280 285 290 295 300 305
 TTA GTT GAA GAA GTG GGC ACT TTC ATG TGC GGC GTG GAA TCC TAC OCT CAG OCT GAA
 L V E E V A T F H C A V E S Y P Q P E>
 310 315 320 325 330 335 340 345 350 355 360 365
 ATT TCT TGG ACC AGA AAT AAA ATT CTC ATC AAG CTG TTT GAC ACC GGC TAC AGC ATC
 I S W T R N K I L I K L F D T R Y S I>
 370 375 380 385 390 395 400 405 410 415 420
 CGA GAG AAC GGT CAG CTC CTC ACC ATC CTG AGT GTG GAG GAC AGT GAT GAT GGC ATC
 R E N G Q L L T I L S V E D S D D G I>
 425 430 435 440 445 450 455 460 465 470 475
 TAC TGC TGC ACA GGC AAC AAT GGA GTG GGA GGA GGC GTG GAA AGT TGT GGC GGC CTG
 Y C C T A N N G V G G A V E S C G A L>
 480 485 490 495 500 505 510 515 520 525 530 535
 CAA GTG AAG ATG AAG OCT AAA ATA ACT OGT OCT CCC ATC AAT GTA AAA ATA ATT GAG
 Q V K H K P K I T R P P I N V K I I E>
 540 545 550 555 560 565 570 575 580 585 590
 GGA TTG AAA GCA GTC CTA CCG TGC ACT ACG ATG GGT AAC CCC AAG CCA TCC GTG TCC
 G L K A V L P C T T H G N P K P S V S>
 595 600 605 610 615 620 625 630 635 640 645 650
 TGG ATT AAG GGG GAC AGT GCT CTC AGG GAA AAT TCC AGG ATT GCA GTT CTT GAA TCT
 W I K G D S A L R E N S R I A V L E S>
 655 660 665 670 675 680 685 690 695 700 705
 GGG AGT TTA AGG ATC CAT AAT GTG CAA AAG GAA GAC GCA GGA CAG TAC CGA TGT GTG
 G S L R I H N V Q K E D A G Q Y R C V>
 710 715 720 725 730 735 740 745 750 755 760
 GCA AAA AAC AGC CTG GGC ACA GCT TAC TCC AAA CTG GTG AAG CTG GAA GTG GAG GTT
 A K N S L G T A Y S K L V K L E V E V>
 765 770 775 780 785 790 795 800 805 810 815 820
 TTT GCA AGA ATC CTG GGT GCT OCT GAA TCC CAC AAT GTC ACC TTT GGT TCC TTT GTA
 F A R I L R A P E S H N V T F G S F V>

Title: NOVEL TYROSINE RECEPTORS
AND LIGANDS

Inventor: Valenzuela, et al.
ATT. DOCKET NO: REG 195-BZ

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Fig. 1B

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      825      830      835      840      845      850      855      860      865      870      875
      *      *      *      *      *      *      *      *      *      *
AOC CTA CGC TGC ACA GCA ATA GGC ATG OCT GTC COC AOC ATC AGC TGG ATT GAA AAC
T   L   R   C   T   A   I   G   M   P   V   P   T   I   S   W   I   E   N>

880      885      890      895      900      905      910      915      920      925      930      935
      *      *      *      *      *      *      *      *      *      *
GGA AAT GCT GTT TCT TCA GGT TOC ATT CAA GAG AAT GTG AAA GAC CGA GTG ATT GAC
G   N   A   V   S   S   G   S   I   Q   E   N   V   K   D   R   V   I   D>

      940      945      950      955      960      965      970      975      980      985      990
      *      *      *      *      *      *      *      *      *      *
TCA AGA CTC CAG CTC TTT ATC ACA AAG OCA GGA CTC TAC ACA TGC ATA GCT AOC AAT
S   R   L   Q   L   F   I   T   K   P   G   L   Y   T   C   I   A   T   N>

995      1000      1005      1010      1015      1020      1025      1030      1035      1040      1045
      *      *      *      *      *      *      *      *      *      *
AAG CAT GGA GAG AAA TTC AGT AOC GCA AAG GCT GCA GOC ACT GTC AGT ATA GCA GAA
K   H   G   E   K   F   S   T   A   K   A   A   A   T   V   S   I   A   E>

1050      1055      1060      1065      1070      1075      1080      1085      1090      1095      1100      1105
      *      *      *      *      *      *      *      *      *      *
TGG AGC AAA TCA CAG AAA GAA AGC AAA GGC TAC TGT GOC CAG TAC AGA GGG GAG GTG
W   S   K   S   Q   K   E   S   K   G   Y   C   A   Q   Y   R   G   E   V>

      1110      1115      1120      1125      1130      1135      1140      1145      1150      1155      1160
      *      *      *      *      *      *      *      *      *      *
TGT GAT GOC GTC CTG GTG AAA GAC TCT CTT GTC TTC TTC AAC AOC TOC TAT OCC GAC
C   D   A   V   L   V   K   D   S   L   V   F   F   N   T   S   Y   P   D>

1165      1170      1175      1180      1185      1190      1195      1200      1205      1210      1215      1220
      *      *      *      *      *      *      *      *      *      *
OCT GAG GAG GOC CAA GAG CTG CTG ATC CAC ACT GOG TGG AAT GAA CTC AAG GCT GTG
P   E   E   A   Q   E   L   L   I   H   T   A   W   N   E   L   K   A   V>

      1225      1230      1235      1240      1245      1250      1255      1260      1265      1270      1275
      *      *      *      *      *      *      *      *      *      *
AGC OCA CTC TGC OGA OCA GCT GOC GAG GCT CTG CTG TGT AAT CAC CTC TTC CAG GAG
S   P   L   C   R   P   A   A   E   A   L   L   C   N   H   L   F   Q   E>

1280      1285      1290      1295      1300      1305      1310      1315      1320      1325      1330
      *      *      *      *      *      *      *      *      *      *
TGC AGC OCT GGA GTG CTA OCT ACT OCT ATG COC ATT TGC AGA GAG TAC TGC TTG GCA
C   S   P   G   V   L   P   T   P   M   P   I   C   R   E   Y   C   L   A>

1335      1340      1345      1350      1355      1360      1365      1370      1375      1380      1385      1390
      *      *      *      *      *      *      *      *      *      *
GTA AAG GAG CTC TTC TGT GCA AAG GAA TGG CTG GCA ATG GAA GGG AAG AOC CAC CGC
V   K   E   L   F   C   A   K   E   W   L   A   M   E   G   K   T   H   R>

      1395      1400      1405      1410      1415      1420      1425      1430      1435      1440      1445
      *      *      *      *      *      *      *      *      *      *
GGA CTC TAC AGA TOC GGG ATG CAT TTC CTC COG GTC COG GAG TGC AGC AAG CTT OCC
G   L   Y   R   S   G   M   H   F   L   P   V   P   E   C   S   K   L   P>

1450      1455      1460      1465      1470      1475      1480      1485      1490      1495      1500      1505
      *      *      *      *      *      *      *      *      *      *
AGC ATG CAC CAG GAC CCC ACA GGC TGC ACA AGA-CTG COG TAT TTA GAT TAT AAA AAA
S   M   H   Q   D   P   T   A   C   T   R   L   P   Y   L   D   Y   K   K>

      1510      1515      1520      1525      1530      1535      1540      1545      1550      1555      1560
      *      *      *      *      *      *      *      *      *      *
GAA AAC ATA ACA ACA TTC COG TOC ATA ACG TOC TOC AAG COG AGC GTG GAC ATT OCA
E   N   I   T   T   F   P   S   I   T   S   S   K   P   S   V   D   I   P>

1565      1570      1575      1580      1585      1590      1595      1600      1605      1610      1615
      *      *      *      *      *      *      *      *      *      *
AAC CTG OCT GCC TOC ACG TCT TOC TTC GOC GTC TOG CCT GOG TAC TOC ATG ACT GTC
N   L   P   A   S   T   S   S   F   A   V   S   P   A   Y   S   M   T   V>

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Title: NOVEL TYROSINE RECEPTORS
AND LIGANDS

Inventor: Valenzuela, et al.

ATT. DOCKET NO: REG 195-BZ

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Fig. 1C

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1620 1625 1630 1635 1640 1645 1650 1655 1660 1665 1670 1675
* * * * *
ATC ATC TCC ATC ATG TCC TGC TTT GCG GTG TTT GCT CTC CTC ACC ATC ACT ACT CTC
I I S I M S C F A V F A L L T I T T L>

1680 1685 1690 1695 1700 1705 1710 1715 1720 1725 1730
* * * * *
TAT TGC TGC CGA AGG AGG AGA GAG TGG AAA AAT AAG AAA AGA GAG TOG GCA GCG GTG
Y C C R R R R E W K N K K R E S A A V>

1735 1740 1745 1750 1755 1760 1765 1770 1775 1780 1785 1790
* * * * *
ACC CTC ACC ACA TTG OCT TCC GAG CTC CTG CTG GAC AGG CTG CAT CCC AAC CCC ATG
T L T T L P S E L L L D R L H P N P M>

1795 1800 1805 1810 1815 1820 1825 1830 1835 1840 1845
* * * * *
TAC CAG AGG ATG CCA CTC CTT CTG AAT CCC AAG TTG CTC AGC CTG GAG TAT CCG AGG
Y Q R M P L L L N P K L L S L E Y P R>

1850 1855 1860 1865 1870 1875 1880 1885 1890 1895 1900
* * * * *
AAT AAC ATC GAG TAT GTC AGA GAC ATC GGA GAG GGA GCG TTT GGA AGG GTC TTT CAA
N N I E Y V R D I G E G A F G R V F Q>

1905 1910 1915 1920 1925 1930 1935 1940 1945 1950 1955 1960
* * * * *
GCG AGG GCG CCA GGC TTG CTT OCT TAT GAA CCC TTC ACT ATG GTG GCT GTG AAG ATG
A R A P G L L P Y E P F T M V A V K M>

1965 1970 1975 1980 1985 1990 1995 2000 2005 2010 2015
* * * * *
CTG AAG GAG GAG GCG TCC GCA GAT ATG CAG GCA GAC TTT CAG AGG GAG GCA GCG CTC
L K E E A S A D M Q A D F Q R E A A L>

2020 2025 2030 2035 2040 2045 2050 2055 2060 2065 2070 2075
* * * * *
ATG GCG GAG TTT GAC AAC CCC AAC ATT GTG AAG CTC TTA GGT GTG TGT GCT GTT GGG
M A E F D N P N I V K L L G V C A V G>

2080 2085 2090 2095 2100 2105 2110 2115 2120 2125 2130
* * * * *
AAG CCA ATG TGC CTG CTC TTT GAA TAT ATG GCG TAT GGT GAC CTC AAT GAG TTC CTC
K P M C L L F E Y M A Y G D L N E F L>

2135 2140 2145 2150 2155 2160 2165 2170 2175 2180 2185
* * * * *
CGA AGC ATG TCC OCT CAC ACT GTG TGC AGC CTC AGC CAC AGT GAC CTG TCC ACG AGG
R S M S P H T V C S L S H S D L S T R>

2190 2195 2200 2205 2210 2215 2220 2225 2230 2235 2240 2245
* * * * *
GCT CGG GTG TCC AGC OCT GGT OCT CCA CCC CTG TCT TGT GCG GAA CAG CTC TGT ATT
A R V S S P G P P P L S C A E Q L C I>

2250 2255 2260 2265 2270 2275 2280 2285 2290 2295 2300
* * * * *
GCG AGG CAA GTG GCA GCT GGC ATG GCG TAC CTG TOG GAG CGC AAG TTT GTC CAT CGG
A R Q V A A G M A Y L S E R K F V H R>

2305 2310 2315 2320 2325 2330 2335 2340 2345 2350 2355 2360
* * * * *
GAC TTA GCT ACC AGG AAC TGC CTG GTT GGA GAG AAC ATG GTG GTG AAA ATT GCA GAC
D L A T R N C L V G E N M V V K I A D>

2365 2370 2375 2380 2385 2390 2395 2400 2405 2410 2415
* * * * *
TTT GCG CTC TCT AGG AAC ATC TAC TCC GCA GAC TAC TAC AAA GCT GAT GGA AAC GAT
F G L S R N I Y S A D Y Y K A D G N D>

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Title: NOVEL TYROSINE RECEPTORS
AND LIGANDS

Inventor: Valenzuela, et al.

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Fig. 1D

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2420 2425 2430 2435 2440 2445 2450 2455 2460 2465 2470
*      *      *      *      *      *      *      *
GCT ATA OCT ATC OGC TGG ATG OCA OCC GAG TCT ATC TTC TAC AAC OGC TAC AOC ACG
A I P I R W M P P E S I F Y N R Y T T>

2475 2480 2485 2490 2495 2500 2505 2510 2515 2520 2525 2530
*      *      *      *      *      *      *      *
GAG TCA GAT GTG TGG GCT TAT GGC GTG GTC CTC TGG GAG ATC TTC TCC TAT GGA CTG
E S D V W A Y G V V L W E I F S Y G L>

2535 2540 2545 2550 2555 2560 2565 2570 2575 2580 2585
*      *      *      *      *      *      *      *
CAG OCC TAC TAT GGA ATG GOC CAT GAG GAG GTC ATT TAC TAT GTG AGA GAT GGT AAC
Q P Y Y G M A H E E V I Y Y V R D G N>

2590 2595 2600 2605 2610 2615 2620 2625 2630 2635 2640 2645
*      *      *      *      *      *      *      *
ATC CTT GOC TGC OCT GAG AAC TGT OCC TTG GAA CTG TAC AAC CTT ATG OGC CTA TGT
I L A C P E N C P L E L Y N L H R L C>

2650 2655 2660 2665 2670 2675 2680 2685 2690 2695 2700
*      *      *      *      *      *      *      *
TGG AGC AAG CTG OCT GCA GAC AGA OCC AGC TTC TGC AGT ATC CAC OGC ATC CTG CAG
W S K L P A D R P S F C S I H R I L Q>

2705 2710 2715 2720 2725 2730 2735 2740 2745 2750 2755 2760
*      *      *      *      *      *      *      *
OGC ATG TGC GAG AGA GCA GAG GGA ACG GTA GGC GTC TAA GGTGAOCC TGCTCAACAA
R M C E R A E G T V G V *>

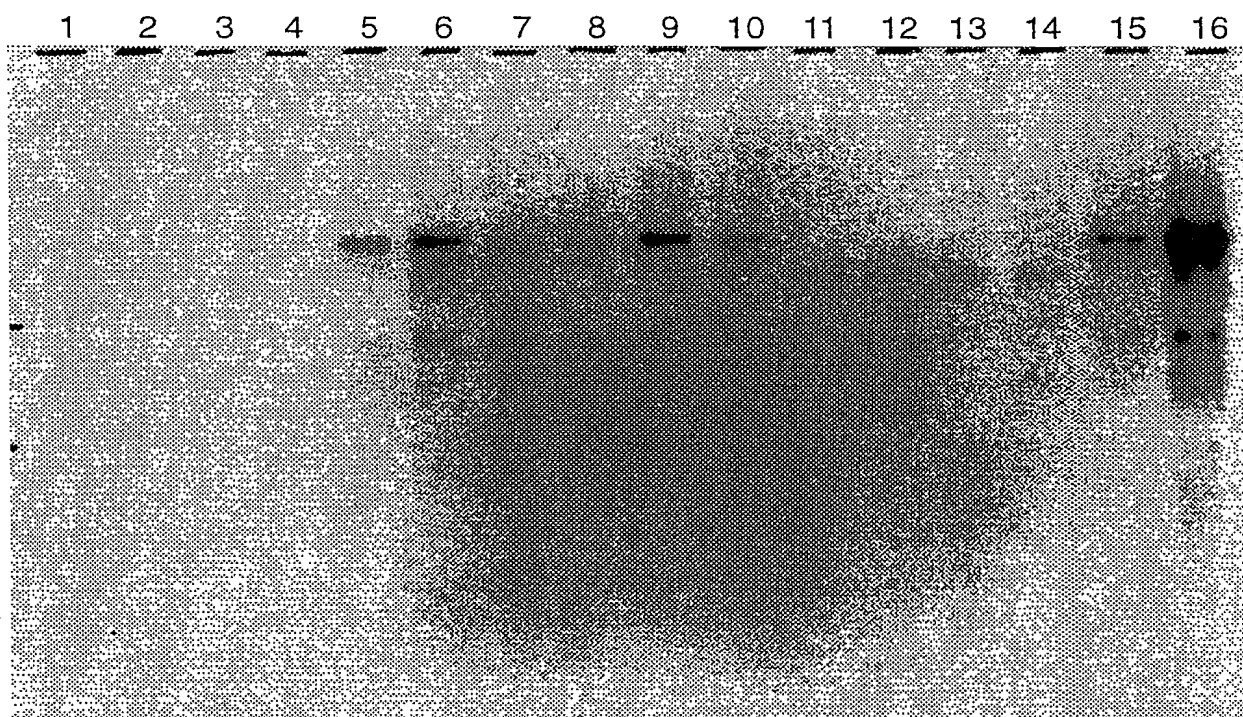
2765 2770 2775 2780 2785 2790 2795 2800 2805 2810 2815 2820 2825 2830
*      *      *      *      *      *      *      *
ACACCCAGGA GGATCTTTTC AGACTGOGAG CTGGAGGGAT OCTAAAGCAG AGGGGOGNATA AGTNCAGATA

2835 2840 2845 2850 2855 2860 2865
*      *      *      *      *
GGAAGAGTTT ATCTCAGGCA GCAOGTNCAG TTGGTTGTT

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Fig.2.

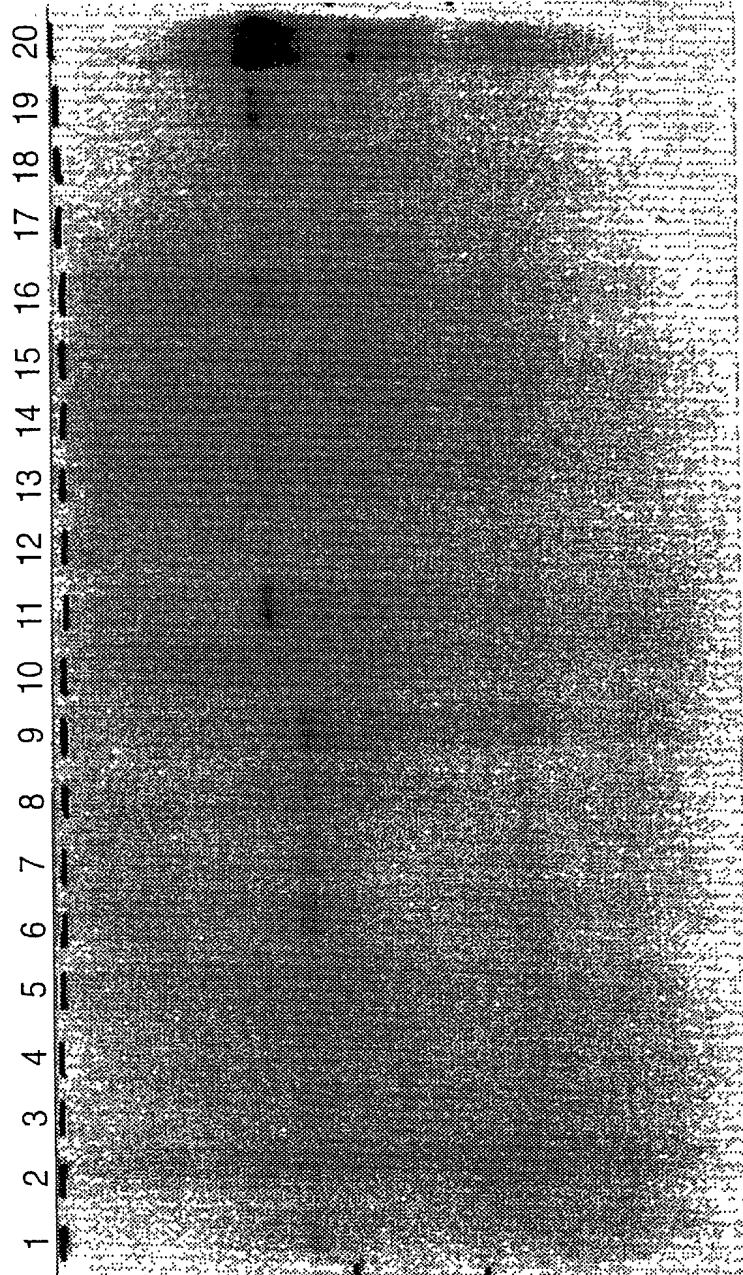


Title: NOVEL TYROSINE RECEPTORS
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Inventor: Valenzuela, et al.
A.I.T. DOCKET NO: REG. 195-BZ

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ATT. DOCKET NO: REG 195-BZ

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Fig. 3.



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Fig. 4A

10	20	30	40	50	60
ATG AGA GAG CTC GTC AAC ATT OCA CTG GTA CAT ATT CTT ACT CTG GTT GGC TTC AOC GGA					
H R E L V N I P L V H I L T L V A F S Q					
70	80	90	100	110	120
ACT GAG AAA CTT OCA AAA GCT OCT GTC ATC AOC ACT OCT CTT GAA ACA GTG GAT GOC TTA					
T E K L P K A P V I T T P L E T V D A L					
130	140	150	160	170	180
GTT GAA GAA GTG GCT ACT TTC ATG TGT GCA GTG GAA TCC TAC CCC CAG OCT GAG ATT TCC					
V E E V A T F H C A V E S Y P Q P E I S					
190	200	210	220	230	240
TGG ACT AGA AAT AAA ATT CTC ATT AAA CTC TTT GAC AOC CCC TAC AGC ATC CCG GAG AAT					
W T R N K I L I K L F D T R Y S I R E N					
250	260	270	280	290	300
GGG CAG CTC CTC ACC ATC CTG AGT GTG GAA GAC AGT GAT GAT GGC ATT TAC TGC TGC ACG					
G Q L L T I L S V E D S D D G I Y C C T					
310	320	330	340	350	360
CCC AAC AAT GGT GTG GGA GGA GCT GTG GAG AGT TGT GGA GGC CTG CAA GTG AAG ATG AAA					
A N N G V G G A V E S C G A L Q V K H K					
370	380	390	400	410	420
OCT AAA ATA ACT CGC CCT CCC ATA AAT GTG AAA ATA ATA GAG GGA TTA AAA GCA GTC CTA					
P K I T R P P I N V K I I E G L K A V L					
430	440	450	460	470	480
CCA TGT ACT ACA ATG GGT AAT CCC AAA CCA TCA GTG TCT TGG ATA AAG GGA GAC AGC CCT					
P C T T H G N P K P S V S W I K G D S P					
490	500	510	520	530	540
CTC AGG GAA AAT TCC CGA ATT GCA GTT CTT GAA TCT GGG AGC TTG AGG ATT CAT AAC GTA					
L R E N S R I A V L E S G S L R I H N V					
550	560	570	580	590	600
CAA AAG GAA GAT GCA GGA CAG TAT CGA TGT GTG GCA AAA AAC AGC CTC GGG ACA GCA TAT					
Q K E D A G Q Y R C V A K N S L G T A Y					
610	620	630	640	650	660
TCC AAA GTG GTG AAG CTG GAA GTT GAG GTT TTT GGC AGG ATC CTG CCG GCT CCT GAA TCC					
S K V V K L E V E V F A R I L R A P E S					
670	680	690	700	710	720
CAC AAT GTC ACC TTT GGC TCC TTT GTG ACC CTG CAC TGT ACA GCA ACA GGC ATT CCT GTC					
H N V T F G S F V T L H C T A T G I P V					

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Fig. 4B

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      730      740      750      760      770      780
      *      *      *      *      *      *
OCC ACC ATC ACC TGG ATT GAA AAC GGA AAT GCT GTT TCT TCT GGG TCC ATT CAA GAG AGT
P   T   I   T   W   I   E   N   G   N   A   V   S   S   G   S   I   Q   E   S

      790      800      810      820      830      840
      *      *      *      *      *      *
GTG AAA GAC CGA GTG ATT GAC TCA AGA CTG CAG CTG TTT ATC AOC AAG CCA GGA CTC TAC
V   K   D   R   V   I   D   S   R   L   Q   L   F   I   T   K   P   G   L   Y

      850      860      870      880      890      900
      *      *      *      *      *      *
ACA TGC ATA GCT ACC AAT AAG CAT GGG GAG AAG TTC AGT ACT GCC AAG GCT GCA GCC ACC
T   C   I   A   T   N   K   H   G   E   K   F   S   T   A   K   A   A   A   T

      910      920      930      940      950      960
      *      *      *      *      *      *
ATC AGC ATA GCA GAA TGG AGT AAA CCA CAG AAA GAT AAC AAA GGC TAC TGC GCC CAG TAC
I   S   I   A   E   W   S   K   P   Q   K   D   N   K   G   Y   C   A   Q   Y

      970      980      990      1000      1010      1020
      *      *      *      *      *      *
AGA GGG GAG GTG TGT AAT GCA GTC CTG GCA AAA GAT GCT CTT GTT TTT CTC AAC ACC TCC
R   G   E   V   C   N   A   V   L   A   K   D   A   L   V   F   L   N   T   S

      1030      1040      1050      1060      1070      1080
      *      *      *      *      *      *
TAT GCG GAC CCT GAG GAG GCC CAA GAG CTA CTG GTC CAC ACG GCC TGG AAT GAA CTG AAA
Y   A   D   P   E   E   A   Q   E   L   L   V   H   T   A   W   N   E   L   K

      1090      1100      1110      1120      1130      1140
      *      *      *      *      *      *
GTA GTG AGC CCA GTC TGC CGG CCA GCT GCT GAG GCT TTG TTG TGT AAC CAC ATC TTC CAG
V   V   S   P   V   C   R   P   A   A   E   A   L   L   C   N   H   I   F   Q

      1150      1160      1170      1180      1190      1200
      *      *      *      *      *      *
GAG TGC AGT CCT GGA GTA GTG CCT ACT CCT ATT CCC ATT TGC AGA GAG TAC TGC TTG GCA
E   C   S   P   G   V   V   P   T   P   I   P   I   C   R   E   Y   C   L   A

      1210      1220      1230      1240      1250      1260
      *      *      *      *      *      *
GTA AAG GAG CTC TTC TGC GCA AAA GAA TGG CTG GTA ATG GAA GAG AAG ACC CAC AGA GGA
V   K   E   L   F   C   A   K   E   W   L   V   H   E   E   K   T   H   R   G

      1270      1280      1290      1300      1310      1320
      *      *      *      *      *      *
CTC TAC AGA TCC GAG ATG CAT TTG CTG TCC GTG CCA GAA TGC AGC AAG CTT CCC AGC ATG
L   Y   R   S   E   H   H   L   L   S   V   P   E   C   S   K   L   P   S   H

      1330      1340      1350      1360      1370      1380
      *      *      *      *      *      *
CAT TGG GAC CCC ACG GCC TGT GCC AGA CTG CCA CAT CTA GAT TAT AAC AAA GAA AAC CTA
H   W   D   P   T   A   C   A   R   L   P   H   L   D   Y   N   K   E   N   L

      1390      1400      1410      1420      1430      1440
      *      *      *      *      *      *
AAA ACA TTC CCA CCA ATG ACG TCC TCA AAG CCA AGT GTG GAC ATT CCA AAT CTG CCT TCC
K   T   F   P   P   H   T   S   S   K   P   S   V   D   I   P   N   L   P   S

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AND LIGANDS

Inventor: Valenzuela, et al.

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Fig. 4C

1450	1460	1470	1480	1490	1500
TCC	TCC	TCT	TCT	TCC	TTC
S	S	S	S	S	F
1510	1520	1530	1540	1550	1560
ATG	TCC	AGC	TTT	GCA	ATA
H	S	S	F	A	I
1570	1580	1590	1600	1610	1620
AGA	AAA	CAA	TGG	AAA	AAT
R	K	Q	W	K	N
1630	1640	1650	1660	1670	1680
TCT	GAG	CTC	TTA	CTA	GAT
S	E	L	L	L	D
1690	1700	1710	1720	1730	1740
CTG	AAC	CCC	AAA	TTG	CTC
L	N	P	K	L	L
1750	1760	1770	1780	1790	1800
ATC	GGA	GAG	GGA	GCG	TTT
I	G	E	G	A	F
1810	1820	1830	1840	1850	1860
GAA	CCT	TTC	ACT	ATG	GTG
E	P	F	T	H	V
1870	1880	1890	1900	1910	1920
GCG	GAC	TTT	CAG	AGG	GAG
A	D	F	Q	R	E
1930	1940	1950	1960	1970	1980
CTA	TTA	GGA	GTG	TGT	GCT
L	L	G	V	C	A
1990	2000	2010	2020	2030	2040
GGT	GAC	CTC	AAT	GAG	TTC
G	D	L	N	E	F
2050	2060	2070	2080	2090	2100
AGT	GAC	TTG	TCT	ATG	AGG
S	D	L	S	H	R
2110	2120	2130	2140	2150	2160
GAG	CAG	CTT	TGC	ATT	GCC
E	Q	L	C	I	A

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Fig. 4D

2170 *				2180 *				2190 *				2200 *				2210 *				2220 *			
TTT	GTT	CAC	CGA	GAT	TTA	GCC	AOC	AGG	AAC	TGC	CTG	GTG	GOC	GAG	AAC	ATG	GTG	GTG	AAA				
F	V	H	R	D	L	A	T	R	N	C	L	V	G	E	N	K	V	V	K				
2230 *				2240 *				2250 *				2260 *				2270 *				2280 *			
ATT	GCC	GAC	TTT	GGC	CTC	TCC	AGG	AAC	ATC	TAC	TCA	GCA	GAC	TAC	TAC	AAA	GCT	AAT	GAA				
I	A	D	F	G	L	S	R	N	I	Y	S	A	D	Y	Y	K	A	N	E				
2290 *				2300 *				2310 *				2320 *				2330 *				2340 *			
AAC	GAC	GCT	ATC	CCT	ATC	CGT	TGG	ATG	CCA	CCA	GAG	TCC	ATT	TTT	TAT	AAC	CGC	TAC	ACT				
N	D	A	I	P	I	R	W	H	P	P	E	S	I	F	Y	N	R	Y	T				
2350 *				2360 *				2370 *				2380 *				2390 *				2400 *			
ACA	GAG	TCT	GAT	GTG	TGG	GCC	TAT	GOC	GTG	GTC	CTC	TGG	GAG	ATC	TTC	TCC	TAT	GOC	CTG				
T	E	S	D	V	W	A	Y	G	V	V	L	W	E	I	F	S	Y	G	L				
2410 *				2420 *				2430 *				2440 *				2450 *				2460 *			
CAG	CCC	TAC	TAT	GGG	ATG	GCC	CAT	GAG	GAG	GTC	ATT	TAC	TAC	GTG	CGA	GAT	GGC	AAC	ATC				
Q	P	Y	Y	G	H	A	H	E	E	V	I	Y	Y	V	R	D	G	N	I				
2470 *				2480 *				2490 *				2500 *				2510 *				2520 *			
CTC	TCC	TGC	CCT	GAG	AAC	TGC	CCC	GTG	GAG	CTG	TAC	AAT	CTC	ATG	CGT	CTA	TGT	TGG	AGC				
L	S	C	P	E	N	C	P	V	E	L	Y	N	L	H	R	L	C	W	S				
2530 *				2540 *				2550 *				2560 *				2570 *				2580 *			
AAG	CTG	CCT	GCA	GAC	AGA	CCC	AGT	TTC	ACC	AGT	ATT	CAC	CGA	ATT	CTG	GAA	CGC	ATG	TGT				
K	L	P	A	D	R	P	S	F	T	S	I	H	R	I	L	E	R	K	C				
2590 *				2600 *				2610 *															
GAG	AGG	GCA	GAG	GGA	ACT	GTG	AGT	GTC	TAA														
E	R	A	E	G	T	V	S	V	*														

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Fig.5.

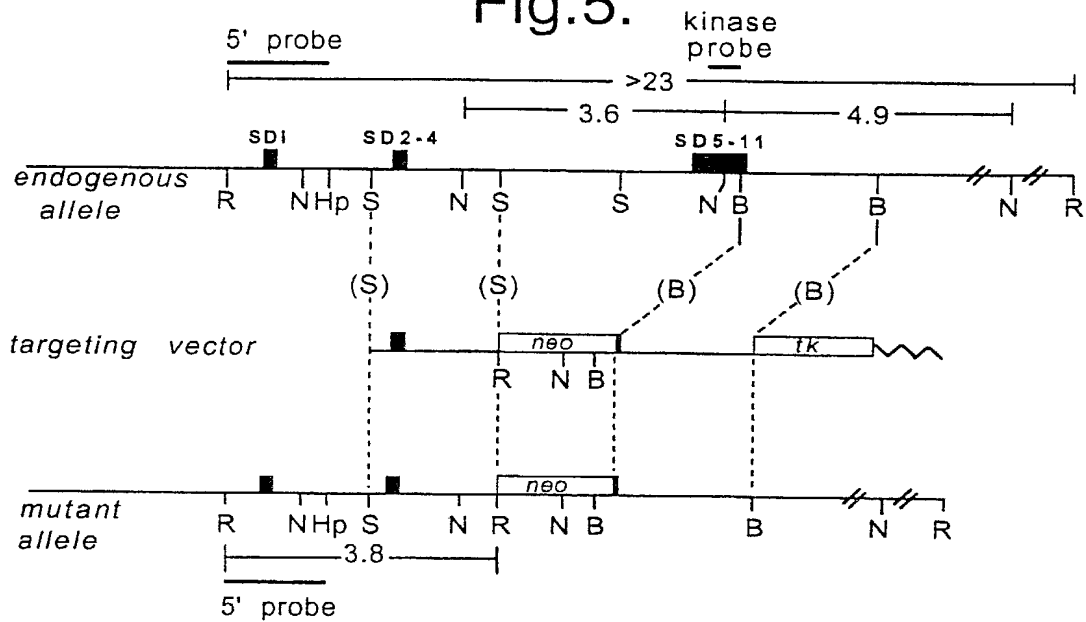


Fig.6.

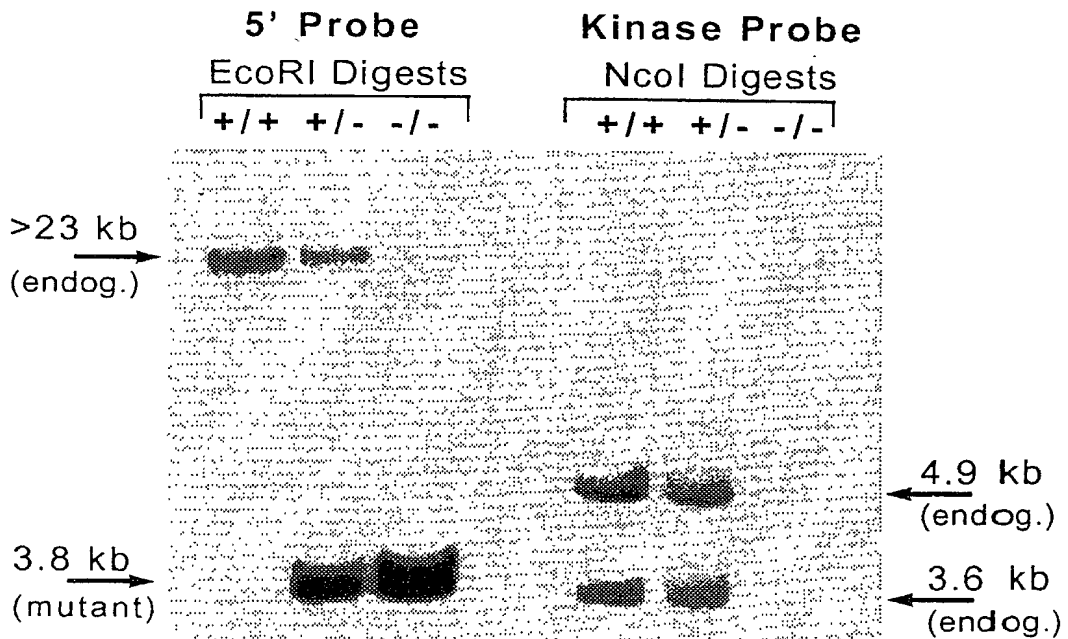


Fig. 7A.

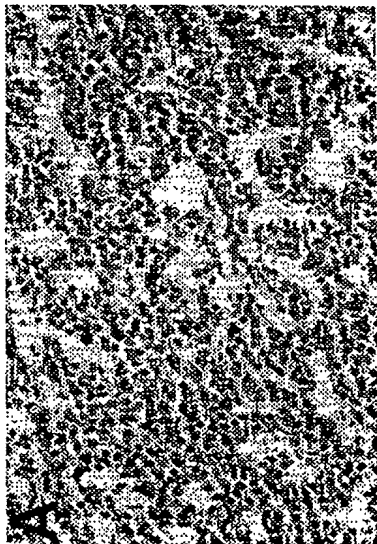


Fig. 7B.

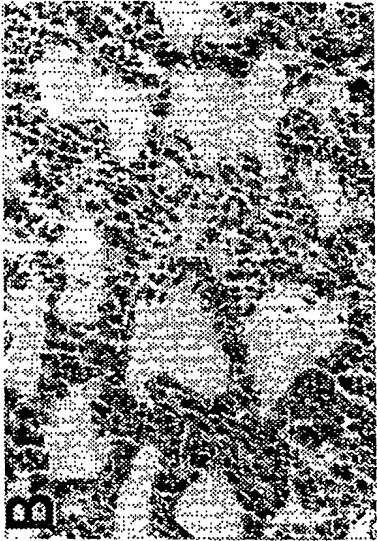


Fig. 7C.

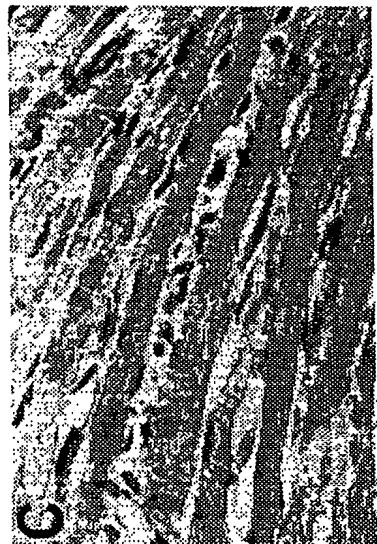


Fig. 7D.



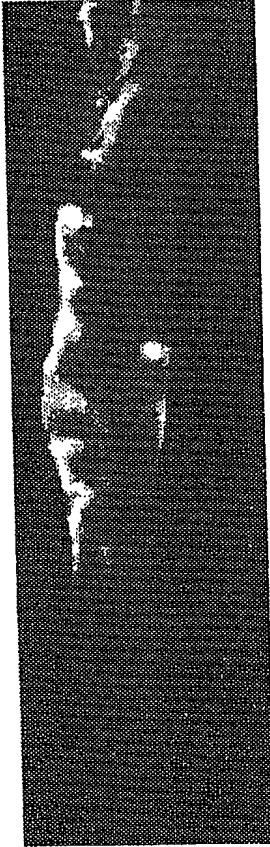
MuSK -/-

CONTROL

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Fig.8A.

Control



MuSK^{-/-}

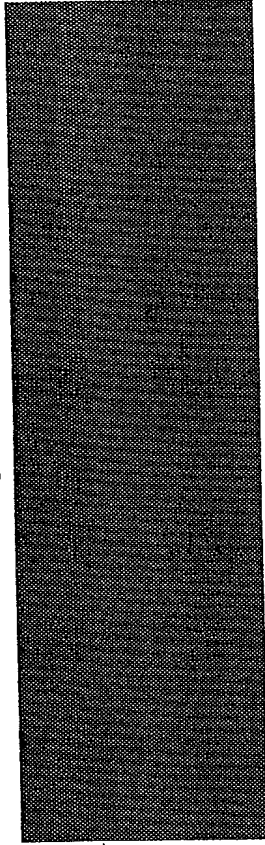


Fig.8B.

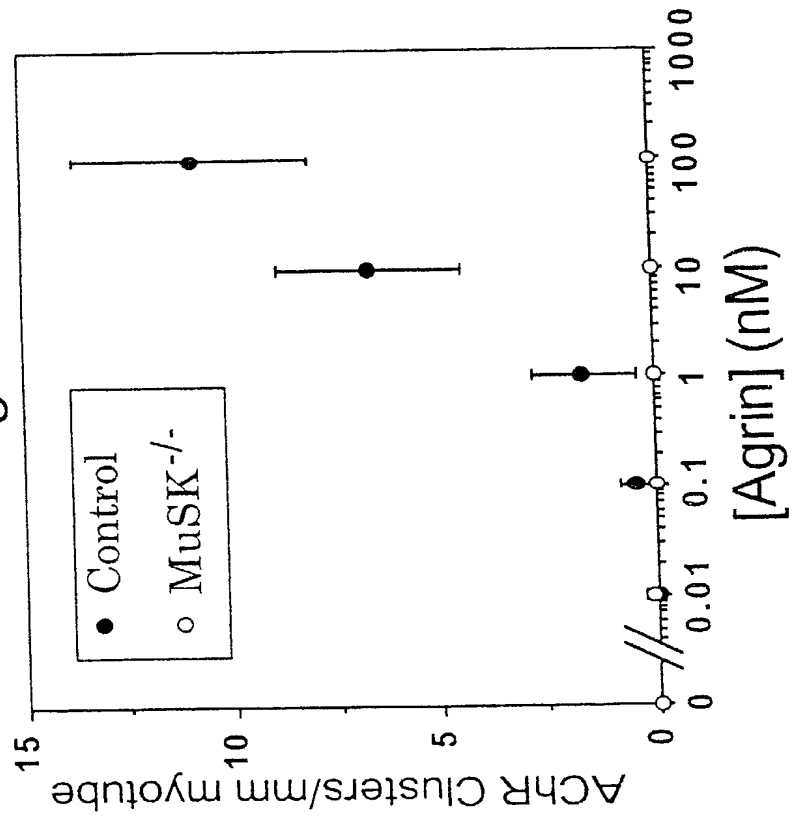
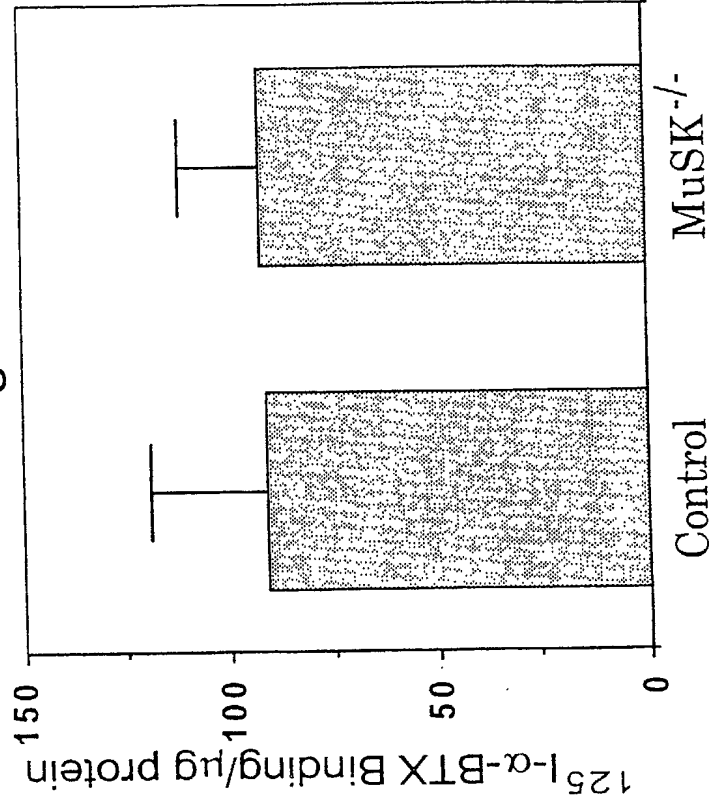


Fig.8C.



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Fig.9A.

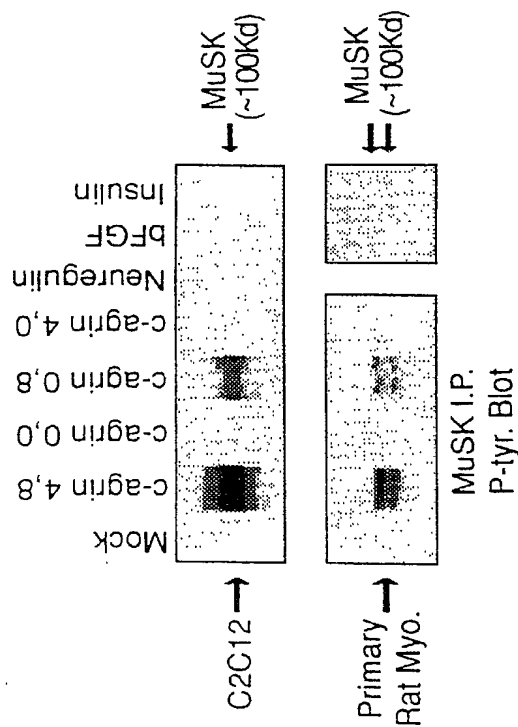


Fig.9B.

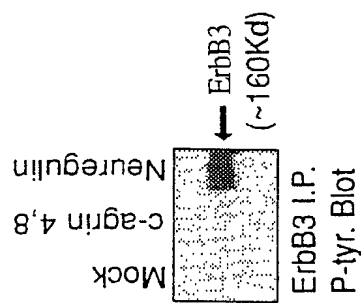


Fig.9C.

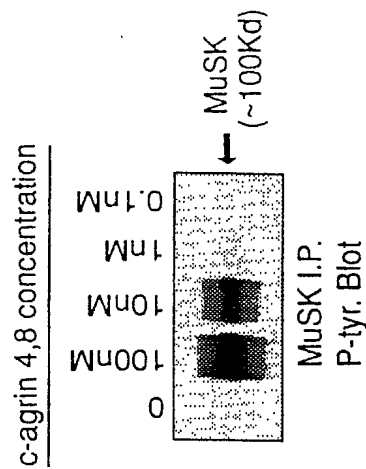


Fig.9D.



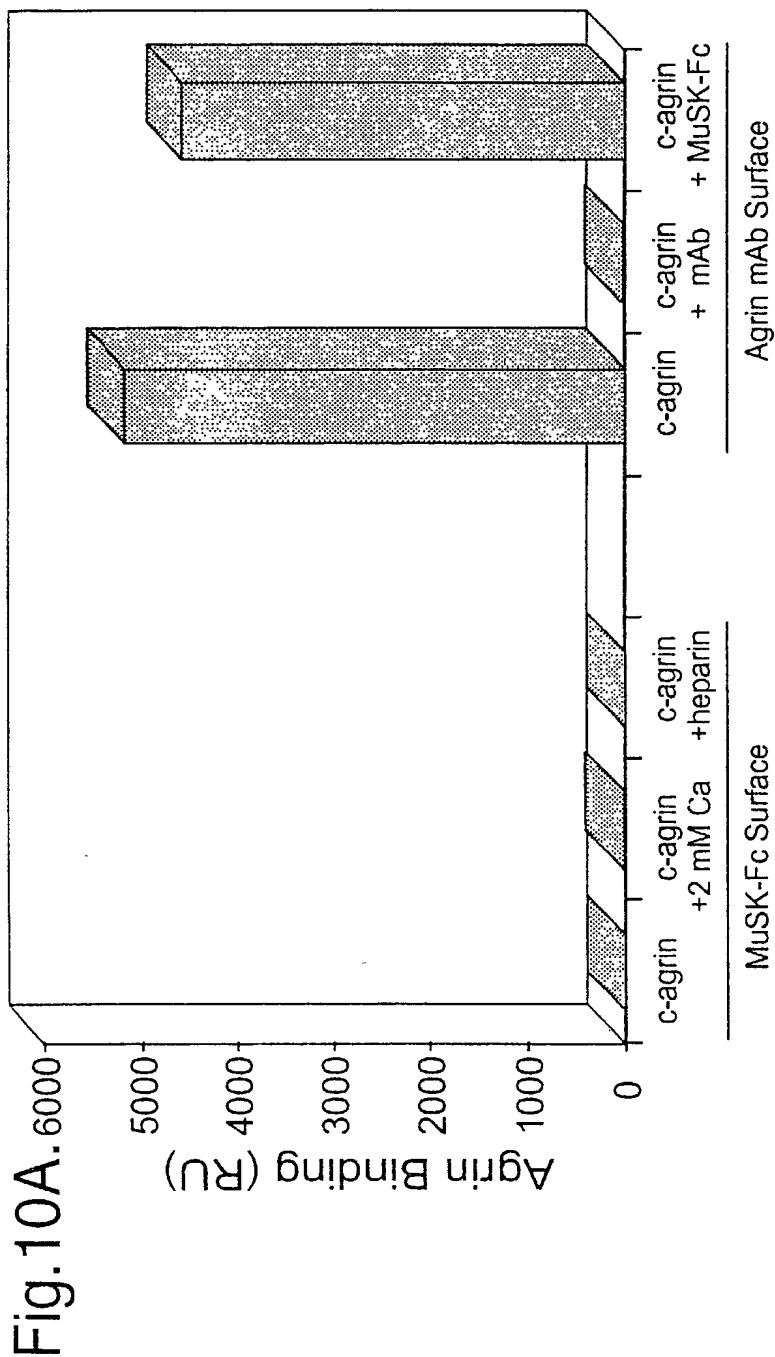
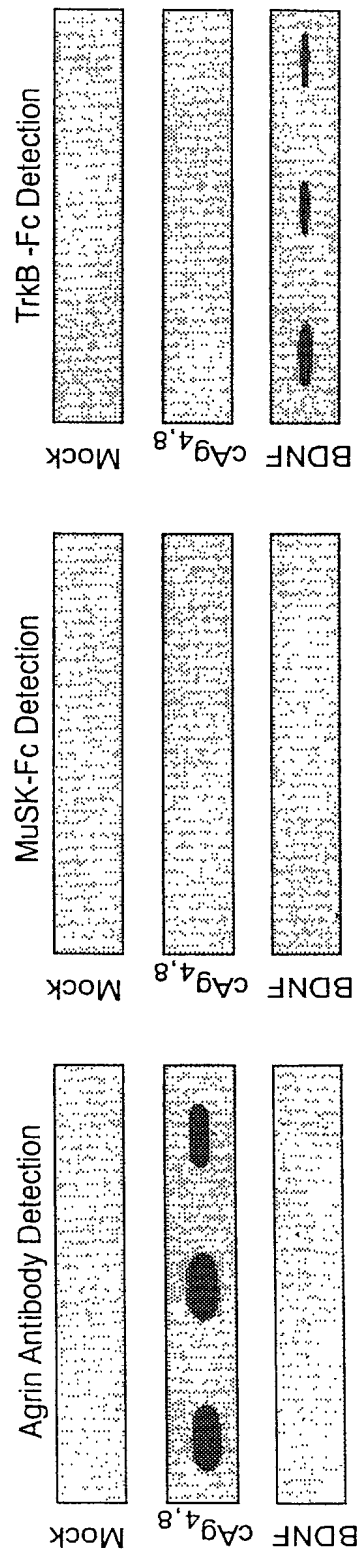
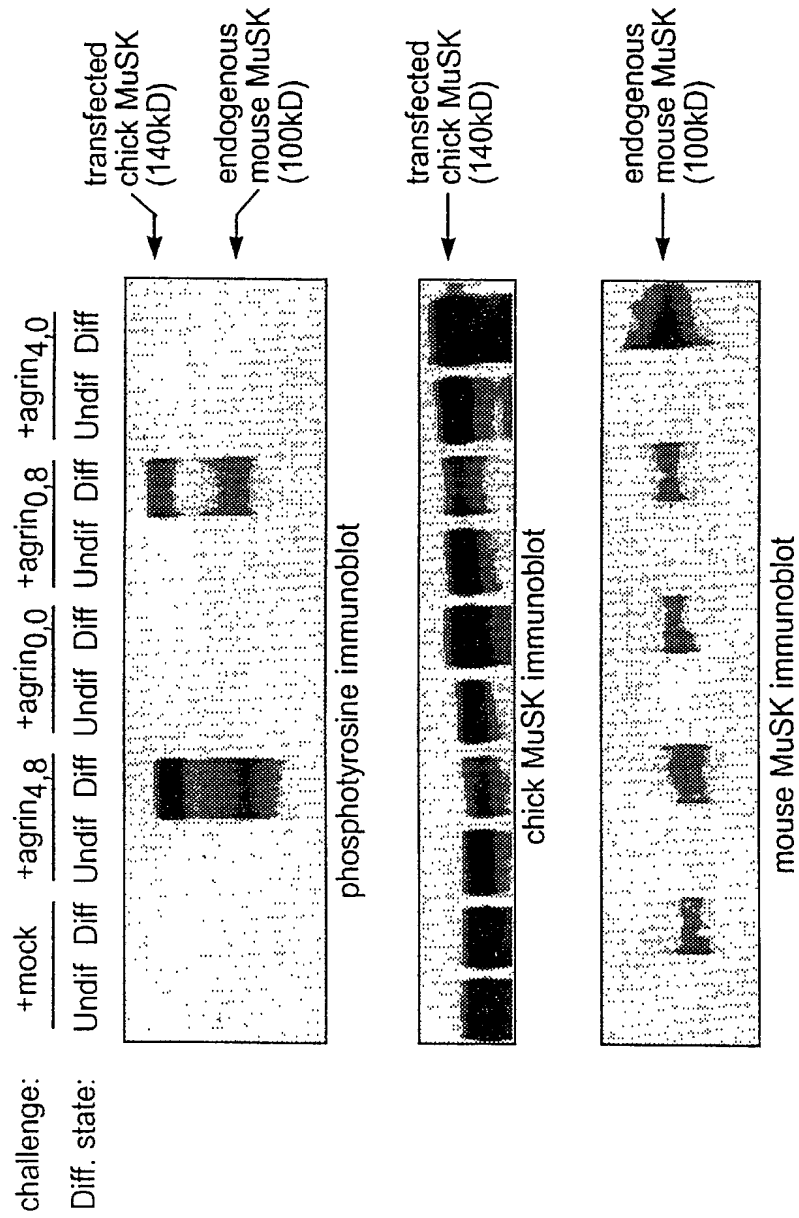


Fig.10B.



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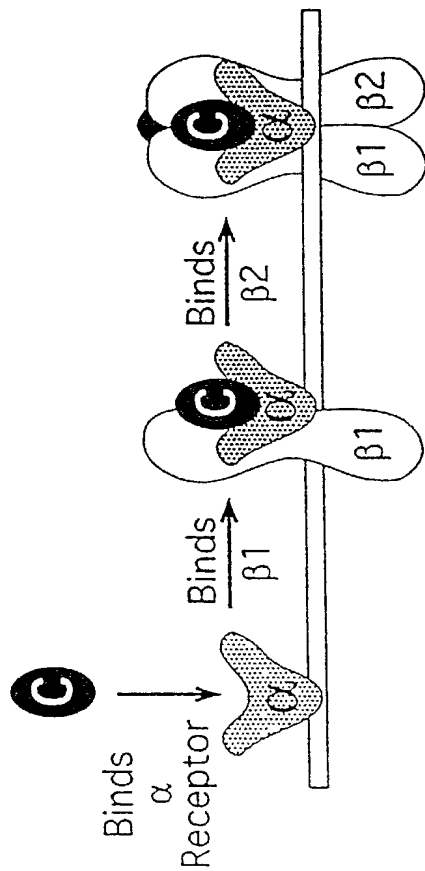
Fig.11.



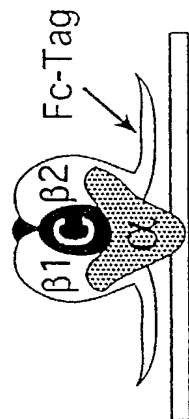
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Fig.12.

A. Formation of CNTF Receptor Complex



B. CNTF Receptor Complex With Soluble β Components



C. Formation of Receptor Complex For Agrin

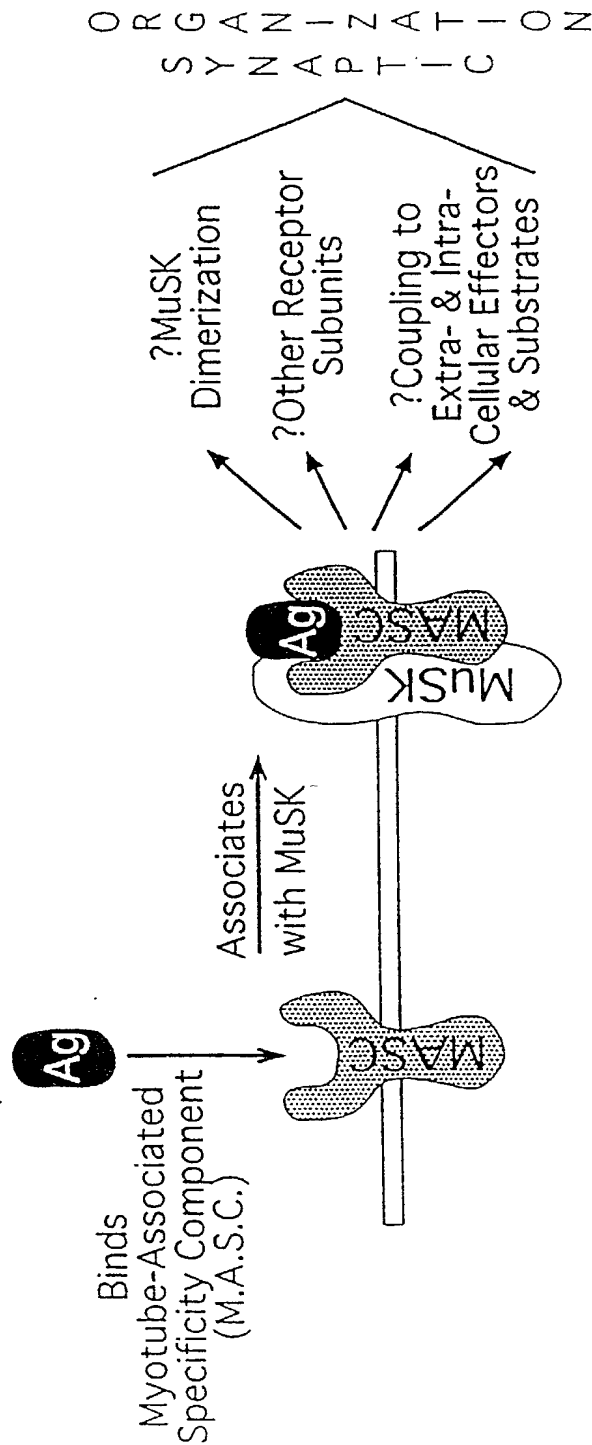


Fig.13A.

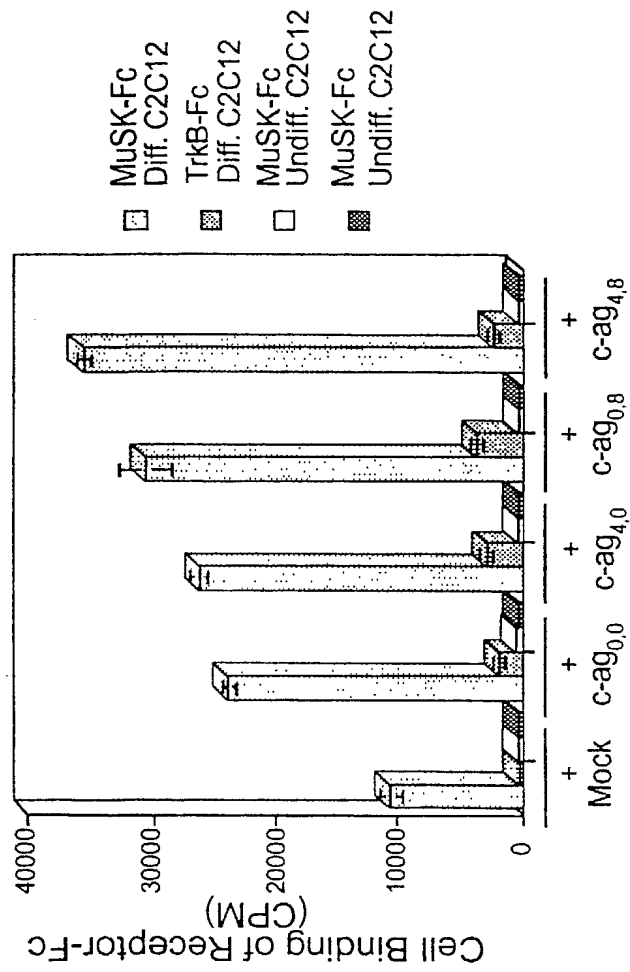


Fig.13B.

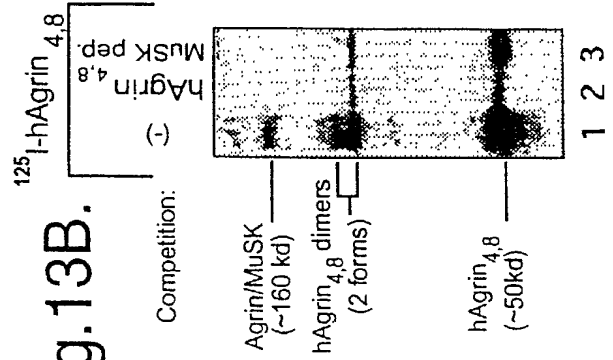
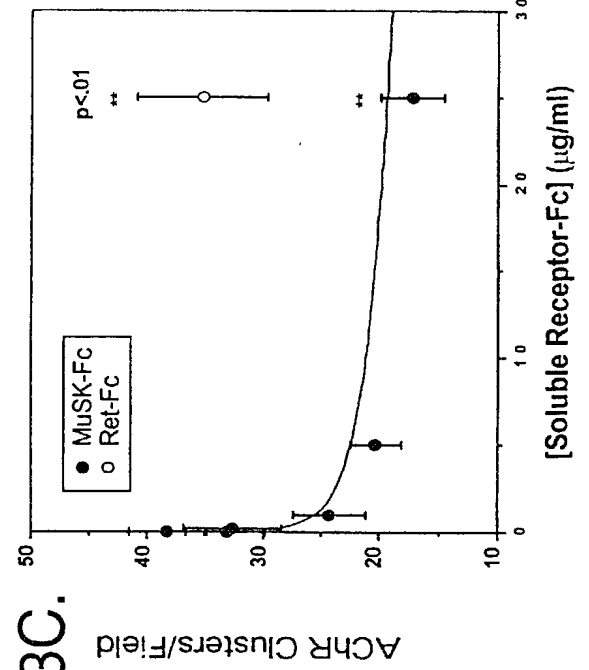


Fig.13C.



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Fig. 14A

10	20	30	40	50	60
* *	* *	* *	* *	* *	* *
MPPLPLEHRP	RQEPGASMLV	RYFMIPCNIC	LILLATSTLG	FAVLLFLSNY	KPGIHFTAP
70	80	90	100	110	120
* *	* *	* *	* *	* *	* *
PTPPDVCRGM	LCGFGAVCEP	SVEDPGRASC	VCKKNACPAT	VAPVCGSDAS	TYSNECELQR
130	140	150	160	170	180
* *	* *	* *	* *	* *	* *
AQCNQRRIR	LLRQGPCGSR	DPCANVTCSF	GSTCVPSADG	QTASCLCPTT	CFGAPDGTVC
190	200	210	220	230	240
* *	* *	* *	* *	* *	* *
GSDGVDYPSE	CQLLSHACAS	QEHIFKKFNG	PCDPCQGSMS	DLNHICRVNP	RTRHPEMLLR
250	260	270	280	290	300
* *	* *	* *	* *	* *	* *
PENCPAQHTP	ICGDDGVTYE	NDCVMSRIGA	TRGLLLQKVR	SGQCQTRDQC	PETCQFNSVC
310	320	330	340	350	360
* *	* *	* *	* *	* *	* *
LSRRGRPHCS	CDRVTCDGSY	RPVCAQDGHT	YNNDWCWRQA	ECRQQRAPP	KHQGPCDQTP
370	380	390	400	410	420
* *	* *	* *	* *	* *	* *
SPCHGVQCAF	GAVCTVKNGK	AECBCQRVCS	GIYDFVCGSD	GVTYGSVCEL	ESMACTLGRE
430	440	450	460	470	480
* *	* *	* *	* *	* *	* *
IQVARRGPCD	PCGQCRFGSL	CEVETGRCVC	PSECVESAQP	VCGSDGHTYA	SECELHVHAC
490	500	510	520	530	540
* *	* *	* *	* *	* *	* *
THQISLYVAS	AGHCQTCGEK	VCTFGAVCSA	GQCVCPRCEH	PPPGPVCSD	GVTYLSACEL
550	560	570	580	590	600
* *	* *	* *	* *	* *	* *
REAACQQQVQ	IEEAHAGPCE	PAECGSGGSG	SGEDDECEQE	LCRQRGGIWD	EDSEDGPCVC
610	620	630	640	650	660
* *	* *	* *	* *	* *	* *
DFSCQSVPRS	PVCGSDGVTY	GTECDLKKAR	CESQQELYVA	AQGACRGPTL	APLLPVAFPH
670	680	690	700	710	720
* *	* *	* *	* *	* *	* *
CAQTPYGCCQ	DNFTAAQGVG	LAGCPSTCHC	NPHGSYSGTC	DPATGQCSCR	PGVGGRLRCR
730	740	750	760	770	780
* *	* *	* *	* *	* *	* *
CEPGFWNFRG	IVTDGHSGCT	PCSCDPRGAV	RDDCEQMTGL	CSCRPGVAGP	KCGQCPDGQV
790	800	810	820	830	840
* *	* *	* *	* *	* *	* *
LHGLGCEADP	MTPVTCVEIH	CEFGASCVEK	AGFAQCICPT	LTCPEANSTK	VCGSDGVTYG

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Fig. 14B

850	860	870	880	890	900
* *	* *	* *	* *	* *	* *
NECQLKAIAC	RQRLDISTQS	LGPCQESVTP	GASPTSASMT	TPRHILSKTL	PFPHNSLPLS
910	920	930	940	950	960
* *	* *	* *	* *	* *	* *
PGSTTHDWPT	PLPISPHTTV	SIPRSTAWPV	LTVPPTAAAS	DVTSLATISIF	SESGSANGSG
970	980	990	1000	1010	1020
* *	* *	* *	* *	* *	* *
DEELSGDEEA	SGGGSGGLEP	FVGSIVVTHG	PPIERASCYN	SPLGCCSDGK	TPSLDSEGSN
1030	1040	1050	1060	1070	1080
* *	* *	* *	* *	* *	* *
CPATKAFQGV	LELEGVEGQE	LFYTPEMADP	KSELFGETAR	SIESTLDDLF	RNSDVKKDFW
1090	1100	1110	1120	1130	1140
* *	* *	* *	* *	* *	* *
SVRLRELPGP	KLVRIVDVH	FDPTTAFQAS	DVGQALLRQI	QVSRPWALAV	RRPLQEHVRF
1150	1160	1170	1180	1190	1200
* *	* *	* *	* *	* *	* *
LDFDWEPTEF	TGAATGTAA	MATARATTVS	RLPASSVTPR	VYPSHTSRPV	GRTTAPPTMR
1210	1220	1230	1240	1250	1260
* *	* *	* *	* *	* *	* *
RPPTTATNMD	RPRTPGHQQP	SKSCDSQPCL	HGGTCQDQDS	GKGFTCSCTA	GRGGSVCEKV
1270	1280	1290	1300	1310	1320
* *	* *	* *	* *	* *	* *
QPPSMPAFKG	HSFLAFPTLR	AYHTLRLALE	FRALETEGLL	LYNGNARGKD	FLALALLDGR
1330	1340	1350	1360	1370	1380
* *	* *	* *	* *	* *	* *
VQFRFDTGSG	PAVLTSLVFV	EPGRWHRLEL	SRHWRQGTLS	VDGETFPVGE	SPSGTDGLNL
1390	1400	1410	1420	1430	1440
* *	* *	* *	* *	* *	* *
DTNLYVGGIP	EEQVAMVLDR	TSVGVGLKGC	IRMLDINNQQ	LELSDWQRAA	VQSSGVGECG
1450	1460	1470	1480	1490	1500
* *	* *	* *	* *	* *	* *
DHPCLPNPCH	GGALCQALEA	GMFLCQCPPG	RFGPTCADEK	SPCQPNPCHG	AAPCRVLSSG
1510	1520	1530	1540	1550	1560
* *	* *	* *	* *	* *	* *
GAKCECPLGR	SGTFCQTVLE	TAGSRPFLAD	FNGFSYLELK	GLHTFERDLG	EKMALEMVFL
1570	1580	1590	1600	1610	1620
* *	* *	* *	* *	* *	* *
ARGPSGLLLY	NGQKTGKGD	FVSLALHNRH	LEFCYDLGKG	AAVIRSKEPI	ALGTWVRVFL
1630	1640	1650	1660	1670	1680
* *	* *	* *	* *	* *	* *
ERNRKGALQ	VGDGPRVLGE	SPKSRKVPHT	MLNLKEPLYI	GGAPDFSKLA	RGAAVSSGFS

▲
Y-site

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Fig. 14C

1690	1700	1710	1720	1730	1740
* *	* *	* *	* *	* *	* *
GVIQLVSLRG	HQLLTQEHVL	RAVDVSPFAD	HPCTQALGNP	CLNGGSCVPR	EATYECLCPG
1750	1760	1770	1780	1790	1800
* *	* *	* *	* *	* *	* *
GFSGLHCEKG	LVEKSVGDL	TLAFDGRTYI	EYLNAVIESE	KALQSNHFEL	SLRTEATQGL
			▲ Z-site		
1810	1820	1830	1840	1850	1860
* *	* *	* *	* *	* *	* *
VLWIGKAAER	ADYMALAI	VDGHLQLSYDLG	SQPVVLRSTV	KVNTNRWLRI	RAHREHREGS
1870	1880	1890	1900	1910	1920
* *	* *	* *	* *	* *	* *
LQVGNEAPVT	GSSPLGATQL	DTDGALWLGG	LQKLPVGQAL	PKAYGTGFVG	CLRDVVVGHR
1930	1940				
* *	* *				
QLHLLDAVT	KPELRPCPTP	*			

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Fig. 15A

ATG	TCT	GCA	CTT	CTG	ATC	CTA	GCT	CTT	GTT	GGA	GCT	GCA	GTT	GCT	GAC
M	S	A	L	L	I	L	A	L	V	G	A	A	V	A	D
TAC	AAA	GAC	GAT	GAC	GAC	AAG	AAG	AGC	CCC	TGC	CAG	CCC	AAC	CCC	TGC
Y	K	D	D	D	D	K	K	S	P	C	Q	P	N	P	C
CAT	GGG	GCG	GCG	CCC	TGC	CGT	GTG	CTG	CCC	GAG	GGT	GGT	GCT	CAG	TGC
H	G	A	A	P	C	R	V	L	P	E	G	G	A	Q	C
GAG	TGC	CCC	CTG	GGG	CGT	GAG	GGC	ACC	TTC	TGC	CAG	ACA	GCC	TCG	GGG
E	C	P	L	G	R	E	G	T	F	C	Q	T	A	S	G
CAG	GAC	GGC	TCT	GGG	CCC	TTC	CTG	GCT	GAC	TTC	AAC	GGC	TTC	TCC	CAC
Q	D	G	S	G	P	F	L	A	D	F	N	G	F	S	H
CTG	GAG	CTG	AGA	GGC	CTG	CAC	ACC	TTT	GCA	CGG	GAC	CTG	GGG	GAG	AAG
L	E	L	R	G	L	H	T	F	A	R	D	L	G	E	K
ATG	GCG	CTG	GAG	GTC	GTG	TTC	CTG	GCA	CGA	GGC	CCC	AGC	GGC	CTC	CTG
M	A	L	E	V	V	F	L	A	R	G	P	S	G	L	L
CTC	TAC	AAC	GGG	CAG	AAG	ACG	GAC	GGC	AAG	GGG	GAC	TTC	GTG	TCG	CTG
L	Y	N	G	Q	K	T	D	G	K	G	D	F	V	S	L
GCA	CTG	CGG	GAC	CGC	CGC	CTG	GAG	TTC	CGC	TAC	GAC	CTG	GGC	AAG	GGG
A	L	R	D	R	R	L	E	F	R	Y	D	L	G	K	G
GCA	GCG	GTC	ATC	AGG	AGC	AGG	GAG	CCA	GTC	ACC	CTG	GGA	GCC	TGG	ACC
A	A	V	I	R	S	R	E	P	V	T	L	G	A	W	T
AGG	GTC	TCA	CTG	GAG	CGA	AAC	GGC	CGC	AAG	GGT	GCC	CTG	CGT	GTG	GGC
R	V	S	L	E	R	N	G	R	K	G	A	L	R	V	G
GAC	GGC	CCC	CGT	GTG	TTG	GGG	GAG	TCC	CCG	AAA	TCC	CGC	AAG	GTT	CCG
D	G	P	R	V	L	G	E	S	P	K	S	R	K	V	P
CAC	ACC	GTC	CTC	AAC	CTG	AAG	GAG	CCG	CTC	TAC	GTA	GGG	GGC	GCT	CCC
H	T	V	L	N	L	K	E	P	L	Y	V	G	G	A	P
GAC	TTC	AGC	AAG	CTG	GCC	CGT	GCT	GCT	GCC	GTG	TCC	TCT	GGC	TTC	GAC
D	F	S	K	L	A	R	A	A	A	V	S	S	G	F	D
GGC	GCC	ATC	CAG	CTG	GTC	TCC	CTC	GGA	GGC	CGC	CAG	CTG	CTG	ACC	CCG
G	A	I	Q	L	V	S	L	G	G	R	Q	L	L	T	P
GAG	CAC	GTG	CTG	CGG	CAG	GTG	GAC	GTC	ACG	TCC	TTT	GCA	GGT	CAC	CCC
E	H	V	L	R	Q	V	D	V	T	S	F	A	G	H	P
TGC	ACC	CGG	GCC	TCA	GGC	CAC	CCC	TGC	CTC	AAT	GGG	GCC	TCC	TGC	GTC
C	T	R	A	S	G	H	P	C	L	N	G	A	S	C	V
CCG	AGG	GAG	GCT	GCC	TAT	GTG	TGC	CTG	TGT	CCC	GGG	GGA	TTC	TCA	GGA
P	R	E	A	A	Y	V	C	L	C	P	G	G	F	S	G
CCG	CAC	TGC	GAG	AAG	GGG	CTG	GTG	GAG	AAG	TCA	GCG	GGG	GAC	GTG	GAT
P	H	C	E	K	G	L	V	E	K	S	A	G	D	V	D

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Fig. 15B

ACC	TTG	GCC	TTT	GAC	GGG	CGG	ACC	TTT	GTC	GAG	TAC	CTC	AAC	GCT	GTG
T	L	A	F	D	G	R	T	F	V	E	Y	L	N	A	V
ACC	GAG	AGC	GAA	CTG	GCC	AAT	GAG	ATC	CCC	GTC	Z-insert				
T	E	S	E	L	A	N	E	I	P	V	E	K	A	L	Q
AGC	AAC	CAC	TTT	GAA	CTG	AGC	CTG	CGC	ACT	GAG	GCC	ACG	CAG	GGG	CTG
S	N	H	F	E	L	S	L	R	T	E	A	T	Q	G	L
GTG	CTC	TGG	AGT	GGC	AAG	GCC	ACG	GAG	CGG	GCA	GAC	TAT	GTG	GCA	CTG
V	L	W	S	G	K	A	T	E	R	A	D	Y	V	A	L
GCC	ATT	GTG	GAC	GGG	CAC	CTG	CAA	CTG	AGC	TAC	AAC	CTG	GGC	TCC	CAG
A	I	V	D	G	H	L	Q	L	S	Y	N	L	G	S	Q
CCC	GTG	GTG	CTG	CGT	TCC	ACC	GTG	CCC	GTC	AAC	ACC	AAC	CGC	TGG	TTG
P	V	V	L	R	S	T	V	P	V	N	T	N	R	W	L
CGG	GTC	GTG	GCA	CAT	AGG	GAG	CAG	AGG	GAA	GGT	TCC	CTG	CAG	GTG	GGC
R	V	V	A	H	R	E	Q	R	E	G	S	L	Q	V	G
AAT	GAG	GCC	CCT	GTG	ACC	GGC	TCC	TCC	CCG	CTG	GGC	GCC	ACG	CAG	CTG
N	E	A	P	V	T	G	S	S	P	L	G	A	T	Q	L
GAC	ACT	GAT	GGA	GCC	CTG	TGG	CTT	GGG	GGC	CTG	CCG	GAG	CTG	CCC	GTG
D	T	D	G	A	L	W	L	G	G	L	P	E	L	P	V
GGC	CCA	GCA	CTG	CCC	AAG	GCC	TAC	GGC	ACA	GGC	TTT	GTG	GGC	TGC	TTG
G	P	A	L	P	K	A	Y	G	T	G	F	V	G	C	L
CGG	GAC	GTG	GTG	GTG	GGC	CGG	CAC	CCG	CTG	CAC	CTG	CTG	GAG	GAC	GCC
R	D	V	V	V	G	R	H	P	L	H	L	L	E	D	A
GTC	ACC	AAG	CCA	GAG	CTG	CGG	CCC	TGC	CCC	ACC	CCA	TGA			
V	T	K	P	E	L	R	P	C	P	T	P	*			

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Fig.16.

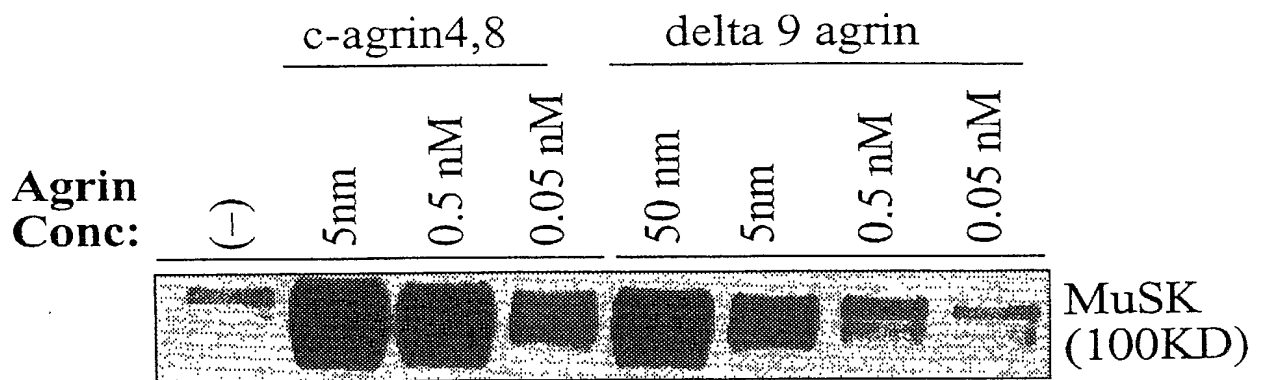


Fig.17.

